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(21) International Application Number: <b>PCT/US98/06371</b>		(72) Inventors; and	
(22) International Filing Date: 1 April 1998 (01.04.98)		(75) Inventors/Applicants (for US only): KLEANTHOS, Harold [GB/US]; 89 Madison Avenue, Newtonville, MA 02160 (US). AL-GARAWI, Amal [SA/US]; 32 Garrison Street #4501, Boston, MA 02114 (US). MILLER, Charles [US/US]; 32 Maple Avenue, Medford, MA 02155 (US). TOMB, Jean-François [-/US]; 3501 St. Paul Street, Baltimore, MD 21222 (US). OOMEN, Raymond, Peter [CA/CA]; R.R. #1, 5400 Lloydtown-Aurora Sideroad, Schomberg, Ontario L0G 1T0 (CA).	
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08/902,615	29 July 1997 (29.07.97)	US	
(63) Related by Continuation (CON) or Continuation-in-Part (CIP) to Earlier Applications		(74) Agent: CLARK, Paul, T.; Clark & Elbing LLP, 176 Federal Street, Boston, MA 02110-2214 (US).	
US	08/833,457 (CON)		
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(54) Title: IDENTIFICATION OF POLYNUCLEOTIDES ENCODING NOVEL <i>HELICOBACTER</i> POLYPEPTIDES IN THE <i>HELICOBACTER</i> GENOME			
(57) Abstract			
The invention provides <i>Helicobacter</i> polypeptides that can be used in vaccination methods for preventing or treating <i>Helicobacter</i> infection, and polynucleotides that encode these polypeptides. The invention also provides diagnostic methods employing these polypeptides.			

CC complexity of the parasitic lifecycle, and provide new targets for  
 CC vaccine and drug development. Parasite resistance to drugs and mosquito  
 CC resistance to insecticides have led to a resurgence of malaria in many  
 CC parts of the world, and there is a pressing need for vaccines and new  
 CC drugs. AA70078 to AA70287 and AAB18144 to AAB18352 represent nucleotide  
 CC and protein sequences given in the present invention, but which are not  
 CC specifically mentioned within the specification.

XX Sequence 1245 AA;

Query Match 13.2%; Score 69; DB 21; Length 1245;  
 Best Local Similarity 27.8%; Pred. NO. 16;  
 Matches 25; Conservative 12; Mismatches 33; Indels 20; Gaps 3;

QY 22 SNKNTVLSKKKPPHLYCVIYIPVLPKLIILFDIAFIPKSLISFONNHVTHNH---T 78  
 DB 1069 SNNYKFFIIRKKKKLKLVCYIMKSFPHI-----LDFWNLSCONEIKNIYKNLHFWIS 1124  
 QY 79 NHNTNIRFNIIIS-----NCRT 95  
 DB 1125 LHNSSIIDFKIINHFLNKIFENISINCTT 1154

RESULT 10.

AAW98706  
 ID AAW98706 standard; Protein; 346 AA.

XX AAW98706;

XX 31-MAR-1999 (first entry)

XX H. Pylori GHPO 727 protein.

XX GHPO protein; Helicobacter infection; gastroduodenal disease; gastritis;  
 KW peptic ulcer disease.

XX Helicobacter pylori.

XX WO9843478-A1.

XX 08-OCT-1998.

XX 01-APR-1998; 98WO-US06371.

XX 29-JUL-1997; 97US-0902615.

XX 01-APR-1997; 97US-0833457.

XX 24-JUN-1997; 97US-0881227.

XX (HUMA-) HUMAN GENOME SCI INC.

XX (INNR) MERIEUX ORAVAX PASTEUR MERIEUX SERUMS.

XX Al-Garawi A, Kleanthous H, Miller C, Oomen RP, Tomb J;  
 DR WPI; 1998-542293/46.

DR N-PSDB; AAX14425.

PT New isolated Helicobacter polynucleotides - used to develop products  
 PT for the diagnosis, prevention and treatment of Helicobacter  
 PT infections and gastrointestinal diseases

PS Claim 8; Page 1482-1484; 2054pp; English.

XX This sequence represents a Helicobacter pylori GHPO protein of the  
 CC invention. The polypeptides can be used for preventing or treating  
 CC Helicobacter infections, and gastroduodenal diseases associated with  
 CC these infections, including acute, chronic, and atrophic gastritis, and  
 CC peptic ulcer diseases, e.g. gastric and duodenal ulcers. They can also be  
 CC used for the production of antibodies. The products can also be used for  
 CC detection and diagnosis.

XX Sequence 346 AA;

XX Match

13.0%; Score 68; DB 19; Length 346;

Best Local Similarity 26.3%; Pred. NO. 4.7;  
 Matches 26; Conservative 12; Mismatches 45; Indels 16; Gaps 3;  
 QY 11 HFKV-----TFWETDLSNNKTLVSLKKKPPHLYCVIYIPVLPKLIIL-----FLDIAF 59  
 DB 146 HFKIKSVIVTYQAVSGAGNKIESLKNELKTALECLEKDPIDLNQVLOAGAFAYPIAF 205  
 QY 60 IPKSLISOFONNHVTHN-----HTNNTNIRFNIIISNC 93  
 DB 206 NATAHIDTFKENCYTKELKMLHETHKIMGVDFPISATC 244

RESULT 11

ABB49049

ID ABB49049 standard; Protein; 907 AA.

XX ABB49049;

XX 05-FEB-2002 (first entry)

XX Listeria monocytogenes protein #1753.

XX Antibacterial; gene therapy; vaccine; biosynthesis; biodegradation;  
 KW vitamin B12; bacterial infection; disease.

XX Listeria monocytogenes.

XX WO200177335-A2.

XX 18-OCT-2001.

XX 11-APR-2001; 2001WO-FR01118.

XX 11-APR-2000; 2000FR-0004629.

XX (INSP) INST PASTEUR.

XX Buchrieser C, Frangeul L, Couve E, Rusniok C, Fsihi H, Dehoux P;  
 PI Dussurget O, Chatouani F, Nedjari H, Glaser P, Kunst F, Cossart P;  
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 PI Perez-Diaz J, Baquero F, Garcia Del Portillo F, Gomez-Lopez N;  
 PI Maduenio E, De Pablos B, Wehland J, Kaerst U, Entian K, Hauf J;  
 PI Rose M, Voss H;

XX WPI; 2002-010914/01.

XX Genomic sequence for Listeria monocytogenes, useful e.g. for treatment  
 XX and prevention of Listeria and related bacterial infections, and  
 XX related polypeptides

XX Claim 6; SEQ ID NO 1754; 192pp; French.

XX The present invention relates to the genome sequence of Listeria  
 CC monocytogenes EGD-e (see ABA03041). The genome sequence and fragments of  
 CC it are useful for selecting probes and primers for detecting genes in L.  
 CC monocytogenes and related organisms, and for studying genetic  
 CC polymorphisms and other genomes. The present sequence is a protein  
 CC encoded by the genome sequence of the present invention. Proteins  
 CC expressed from the genome sequence are useful for raising specific  
 CC antibodies, identification of L. monocytogenes and related organisms,  
 CC for biosynthesis and biodegradation, especially biosynthesis of Vitamin  
 CC B12. The genome sequence and proteins encoded by it are also useful for  
 CC selecting compounds that regulate gene expression and cell replication  
 CC and modulate L. monocytogenes-related diseases. In addition, the genome  
 CC sequence and proteins encoded by it are useful in pharmaceutical and  
 CC vaccines compositions for the treatment or prevention of infections by L.  
 CC monocytogenes and related organisms.  
 CC Note: The sequence data for this patent did not form part of the printed  
 CC specification, but was obtained in electronic format directly from WIPO  
 CC at ftp.wipo.int/pub/published\_pct\_sequences.